

SEQUENCE LISTING

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<110> Williams, Richard B.
 <120> SYSTEM AND METHODS FOR NUCLEIC ACID AND
   POLYPEPTIDE SELECTION
 <130> PRONOV.001BPC
 <150> 09/859,809
 <151> 2001-05-17
<150> 60/206,016
<151> 2000-05-19
 <150> 60/346,965
 <151> 2001-11-16
 <150> 60/529,331
 <151> 2003-12-12
: <150> PCT/US02/37103
 <151> 2002-11-18.
 <150> 10/847,087
 <151> 2004-05-17
 <150> 60/625,707
 <151> 2004-11-05
 <150> 10/847,484
 <151> 2004-05-17
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    · chemically synthesized fragment
 <221> modified base
 <222> (6)...(6)
 <223> n=p
 <400> 1
                                                                      13
 cuagancugg agg
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 <223> Furan sided psoralentated RNA fragment; chemically
       synthesized
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 <222> (2)...(3)
 <223> psoralen bound to UA
 <221> modified base
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cuagancugg agg
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<222> (1)...(5)
<223> n=g, a, u, or c
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nnnnnccucc agaucuagnn nnn
                                                                    23
<210> 4
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<223' Chemically synthesized psoralentated RNA fragment
<221> misc feature
<222> (16)...(17)
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<222> (1) ... (5)
<223> n=g, a, u, or c
<221> misc feature
<222> (19)...(23)
<223> n=g, \bar{a}, u, or c
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<400> 4
  nnnnnccucc agaucuagnn nnn
                                                                      23
  <210> 5
 <211> 13
 <212> RNA
 <213> Artificial Sequence
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 <223> Chemically synthesized fragment 2; psoralentated
       RNA fragment
 <221> modified base
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 <223> psoralen bound to G
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 cuagancugg agg
                                                                     13
 <210> 6.
 <211> 30
 <212> DNA
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<223> chemically synthesized fragment 3; modified tRNA;
       thymine at residue 8 before pseudouridine
<221> modified_base
<222> (9)...(9)
<223> n=p
<221> misc_feature
<222> (30)...(30)
<223> n=puromycin
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uccugugtnc gauccacaga auucgcaccn
                                                                    30
<210> 7
<211> 43
<212> DNA
<213> Artificial Sequence
<223> Chemically synthesized fragment 2+3; modified
      tRNA; thymine at residue 21 before pseudouridine
<221> modified base
<222> (22)...(22)
<223> n=p
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<221> misc_feature
<222> (43)...(43)
<223> n=puromycin
<221> modified base
<222> (6)...(6)
<223> n=p
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cuagancugg agguccugug tncgauccac agaauucgca ccn
                                                                       43
<210> 8
<211> 33
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<400> 8
geggauuuag cucaguuggg agagegeeag acu
                                                                       33
<210> 9
<21.1> 76..
<212> DNA
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<223> Chemically synthesized fragment 1 + 2 + 3;
      modified tRNA; thymine at residue 54 before
      pseudouridine
<221> modified_base
\langle 222 \rangle \ (39) \dots (\overline{3}9)
<223> n=p
<221> modified base
<222> (55) ... (55)
<223> n=p
<221> misc_feature
<222> (76) ... (76)
<223> n=puromycin
<221> misc feature
<222> (35)...(36)
<223> psoralen binding at UA position
<400> 9
geggauuuag cucaguuggg agagegeeag acucuagane uggaggueeu gugtnegaue 60
cacagaauuc gcaccn
<210> 10
<211> 32
<212> RNA
<213> Artificial Sequence
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 <223> chemically synthesized fragment 1; 3' hydroxyl at
        terminus
  <400> 10
 gcggauuuag cucaguuggg agagcgccag ac
                                                                       32
 <210> 11
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 <212> RNA
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 <223> chemically synthesized fragment 2; psoralentated
 <221> modified base
 <222> (1)...(1)
 <223> N=p
 <221> modified base
 <222> (7)...(7)
 <223> N=p
<400> 11
ncuaacnc '
                                                                      8
<210> 1.2
<211> 36
<212> DNA
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<223> Mixed DNA/RNA chemically synthesized fragment 3
<221> modified_base
<222> (15)...(15)
<223> n=p
<221> misc_feature
<222> (36) ... (36)
<223> n=puromycin
uggagguccu gugtncgauc cacagaauuc gcaccn
                                                                      36
<210> 13
<211> 18
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<221> modified base
\langle 222 \rangle (4) \dots (4)
<223> n=p
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<221> modified base
<222> (11) . . . (12)
<223> n=p
<400> 13
cccnccagag nnagaccc
                                                                       18
<210> 14
<211> 14
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<213> Artificial Sequence
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<221> modified_base
<222> (7)...(7)
<223> n=p
<221> modified base
<222> (9)...(9)
<223> n=p
<400> 14
ucuaagneng gagg
                                                                       14
<210> 15
<211> 73
<212> DNA
<213> Artificial Sequence
<223> chemically synthesized; mixed DNA/RNA; alternate
      psoralentated Fragment 1 + 2 + 3
<221> modified base
<222> (32)...(32)
<223> N-3-methyl uridine
<221> misc_feature
<222> (36)...(37)
<223> psoralen bound to UA
<221> modified base
<222> (40) ... (40)
<223> N=p
<221> modified_base
\langle 222 \rangle \ (42) \dots (\overline{4}2)
<223> N=p
<221> misc_feature
<222> (73) ... (73)
<223> N=puromycin
<400> 15
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gcggauuuag cucaguuggg agagcgccag anuucuaagn cnggaggucc ugugtycgau 60
ccacagaauu cgn
<210> 16
<211> 32
<212> RNA
<213> Artificial Sequence
<220>
<223> Chemically synthesized fragment 1; 3' hydroxyl at
      terminus
<400> 16
gcggauuuag cucaguuggg agagcgccag ac
                                                                     32
<210> 1.7
<211> 8
<212> RNA
<213> Artificial Sequence
<223> Chemically synthesized fragment 2; 3' hydroxyl at
      terminus
<221>_modified_base
<222> (1)...(1)
<223> n=p
<221> modified base
<222> (7)...(7)
<223> n=p
<400> 17
                                                                     8
ncuaaanc
<210> 18
<211> 36
<212> DNA
<213> Artificial Sequence
<220>
<223> Chemically synthesized; mixed DNA/RNA fragment 3
<221> modified base
\langle 222 \rangle (15)...(15)
<223> N=p
<221> misc_feature
<222> (36)...(36)
<223> N=puromycin
<400> 18
                                                                     36
uggagguccu gugtncgauc cacagaauuc gcaccn
<210> 19
<211> 21
<212> RNA
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<213> Artificial Sequence	
<220> <223> Chemically synthesized; modified RNA fragment	
<221> modified_base <222> (9)(11) <223> N=p	
<400> 19 cccccgann nagaccccc c	21
<210> 20 <211> 26 <212> RNA <213> Artificial Sequence	
<220> <223> chemically synthesized; seq A1	
<400> 20	
auauauau auauauauau gggggg	26
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<220> <223> Chemically synthesized; seq A2	
<400> 21 cccccatat atatatat atatat	26
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<400> 22 geggauuuag eucaguuggg agagegecag a	31
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<221> misc_feature
 <222> (1)...(1)
 <223> no phosphorylation on 5' end
 <400> 23
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 <223> Chemically synthesized RNA sequence
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 <223> crosslinker between residue 35 and 36
 <221> misc_feature
 <222> (36) . . . (36)
 <223> phosphorylated
 <400>. 24
 geggauuuag eucaguuggg agagegeeag acueua
                                                                      36
 <210> 25
·<211> 36 ·
<212> RNA " `
<213> Artificial Sequence
<223> Chemically synthesized RNA sequence
<221> misc_feature
<222> (1) ... (1)
<223> hydroxylated
<221> misc feature
<222> (35) . . . (36)
<223> crosslinker between residues 35 and 36
<221> misc_feature
<222> (36) ... (36)
<223> phosphorylated
<400> 25
ggggcuuuag cucaguuggg agagcgccag acucua
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<210> 26
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<212> RNA
<213> Artificial Sequence
<220>
<223> Chemically synthesized RNA sequence
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<221> misc feature
 <222> (59) ... (60)
 <223> crosslinker between residues 59 and 60
 ggguuaacuu uagaaggagg ucgccaccau gguuaaaaug aaaaugaaaa ugaaaaugua 60
 <210> 27
 <211> 55
 <212> RNA
 <213> Artificial Sequence
 <220>
 <223> Chemically synthesized RNA sequence M1
ggguuaacuu uagaaggagg ucgccaccau gguuaaaaug aaaaugaaaa ugaaa
 <210> 28
 <211> 61
 <212> RNA
<213> Artificial Sequence
.<220>
<223> Chemically synthesized RNA sequence
<221> misc_feature
<222> (59) ... (60)
<223> crosslink between residue 59 and 60
<221> misc_feature
<222> (61)...(61)
<223> g bound to biotin
<221> misc feature
<222> (33)...(34)
<223> n=g, a, u, or c
<400> 28
ggguuaacuu uagaaggagg ucgccaccau ggnnaaaaug aaaaugaaaa ugaaaaugua 60
<210> 29
<211> 61
<212> RNA
<213> Artificial Sequence
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<223> Chemically synthesized RNA sequence
ggguuaacuu uagaaggagg ucqccaccau qquuaaaauq aaaauqaaaa uqaaaauqua 60
<210> 30
<211> 21
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<212> RNA
 <213> Artificial Sequence
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 <223> Chemically synthesized RNA sequence; generic
       version of SEQ ID NO: 19
 <221> misc feature
 <222> (1) ... (6)
<223> n=g, a, c, or u
<221> modified base
\langle 222 \rangle \ (9) \dots (1\overline{1})
<223> n=p
<221> misc_feature.
<222> (15)...(21)
<223> n=g, a, c, or u
<400> 30
nnnnnngann nagannnnnn n
                                                                      21
<210> 31
<211>.36
<212> DNA
<213> Artificial Sequence
<223> Chemically synthesized; mixed DNA/RNA sequence
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<222> (1)...(1)
<223> phosphorylated
<221> misc_feature
<222> (14)...(14)
<223> thymine
<221> modified base
<222> (15)...(15)
<223> n=p
<221> misc_feature
<222> (36)...(36)
<223> n=puromycin
<400> 31
uggagguccu gugtncgauc cacagaauuc gcaccn
                                                                      36
<210> 32
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<212> DNA
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<220>
<223> Chemically synthesized; Mixed DNA/ RNA sequence
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<221> misc_feature
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  <223> phosphorylated
  <221> misc feature
  <222> (14)...(14)
  <223> thymine
  <221> modified_base
  <222> (15)...(15)
  <223> n=p
<221> misc_feature
<222> (36)...(36)
  <223> puromycin attached at 3' end
36
  uggagguecu gugtnegaue cacagaaueu ecacea
  <210> 33
  <211> 72
  <212> DNA
<213> Artificial Sequence
<220>
<223> mixed DNA/RNA chemically synthesized primer for
SARS-CoV genome sequence
SARS-CoV genome sequence
 <222> (32)...(32)
 <223> N-3-methyl uridine
 <221> misc feature
  <222> (35)...(36)
 <223> psoralen bound to UA
 <221> modified base
  <222> (39)...(39)
 <223> p
 <221> modified base
 <222> (41)...(41)
 <223> p
 <221> misc_feature
 <222> (72)...(72)
 <223> n=puromycin
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 geggauuuag eucaguuggg agagegeeag anucuaagne nggaggueeu gugtyegaue 60
 cacagaauuc gn
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